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Figure 1A

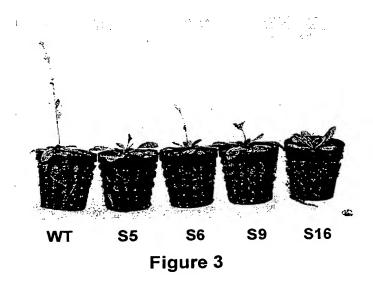
Figure 1B

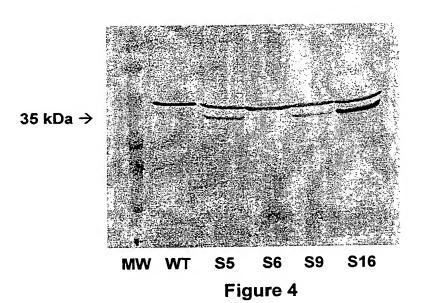


Figure 2A

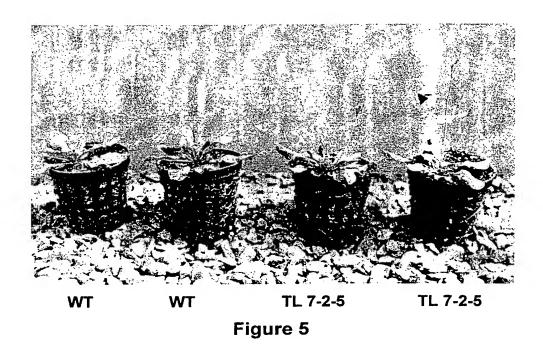
Figure 2B

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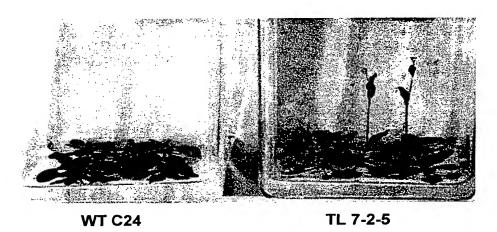


Figure 6

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Figure 7

SEQ ID NO 1:

ATGGCTACCTCAAGCATGAAGAGCATTCCAATGGCGATCCCAAGTTTCTCCATGTGTCAC AAGCTCGAGCTCCTTAAAGAAGGCAAAACTCGCGACGTCCCGAAAGCCGAAGAAGATGAA GGGCTAAGCTGCGAGTTCCAAGAGATGTTGGATTCTCTTCCTAAGGAGAGAGGATGGAGA ACTCGTTACCTTTACCTATTCCAAGGGTTTTGGTGCCAAGCCAAAGAGATTCAAGCCATC ATGTCTTTCCAAAAACATTTCCAATCCCTCGAAAACGACGTCGTTCTCGCCACCATACCT AAATCCGGTACAACCTGGCTAAAAGCTTTAACTTTCACCATCCTTAACCGTCACCGGTTT GATCCGGTTGCCTCGAGTACCAACCACCCTCTTTTCACTTCCAACCCTCATGACCTTGTA CCTTTCTTCGAGTACAAGCTTTACGCCAACGGAGATGTTCCCGATCTCTCGGGTCTAGCC AGTCCAAGAACGTTCGCAACCCACTTACCGTTCGGTTCCCTAAAGGAAACGATCGAGAAA CCCGGTGTGAAGGTCGTGTACTTGTGCCGGAACCCGTTTGACACATTCATCTCTTCGTGG CATTACACCAACAACATCAAATCCGAGTCAGTGAGCCCAGTCTTGCTAGACCAAGCTTTT GATCTGTATTGCCGGGGAGTGATCGGGTTTGGCCCGTTTTGGGAACACATGTTGGGATAC TGGAGAGAGACTTGAAGAGACCAGAGAAAGTCTTCTTTTAAGGTACGAGGATCTCAAA GACGACATCGAGACCAACTTGAAGAGGCTTGCAACTTTCTTAGAGCTTCCTTTCACCGAA GAAGAGGAACGAAAGGGAGTTGTGAAGGCTATCGCCGAGCTGTGTAGCTTCGAGAATCTG AAGAAGTTGGAGGTGAACAAGTCAAACAAGTCGATCAAGAACTTTGAGAATCGATTCTTG TTTCGGAAAGGAGAAGTGAGTGATTGGGTTAACTATTTGTCACCTTCACAAGTGGAAAGA TTGTCAGCCTTAGTGGATGACAAGTTAGGTGGATCTGGTCTCACTTTCAGGTTGAGC

Figure 8

SEQ ID NO 3:

MATSSMKSIPMAIPSFSMCHKLELLKEGKTRDVPKAEEDEGLSCEFQEMLDSLPKERGWR TRYLYLFQGFWCQAKEIQAIMSFQKHFQSLENDVVLATIPKSGTTWLKALTFTILNRHRF DPVASSTNHPLFTSNPHDLVPFFEYKLYANGDVPDLSGLASPRTFATHLPFGSLKETIEK PGVKVVYLCRNPFDTFISSWHYTNNIKSESVSPVLLDQAFDLYCRGVIGFGPFWEHMLGY WRESLKRPEKVFFLRYEDLKDDIETNLKRLATFLELPFTEEEERKGVVKAIAELCSFENL KKLEVNKSNKSIKNFENRFLFRKGEVSDWVNYLSPSQVERLSALVDDKLGGSGLTFRLS WO 01/02589

10019931.05100E 10/019931 PCT/CA00/00801

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Figure 9

SEQ ID NO 2:

GAAGGCCAAGAAGAAGAAGGGCTAAGCTACGAGTTCCAAGAGATGTTGGACTCTCTTCCT AAGGAGAGAGGACGGAGAAATCGTTACCTTTACTTATTCCAAGGGTTTCGGTGCCAAGCT AAGGAGATTCAAGCTATCACGTCTTTCCAAAAACATTTTCAGTCCCTTCCAGACGACGTT GTCCTCGCCACCATACCTAAATCTGGCACAACCTGGTTAAAAGCTTTAACTTTCACCATC CTTACCCGTCATCGGTTTGATCCGGTTTCCTCATCAAGTTCCGACCACCCTCTTCTCACA TCCAACCCTCACGACCTCGTACCTTTCTTCGAGTACAAGCTTTACGCCAACGGAAATGTT CCCGATCTCTCGGGTCTAGCCAGTCCAAGAACATTCGCAACCCACGTACCGTTCGGTGCC CTTAAGGATTCGGTCGAGAATCCCAGTGTGAAGGTTGTGTACCTGTGCCGGAACCCGTTT GACACATTCATCTCCATGTGGCATTACATCAACAACATCACTTCCGAGTCAGTGAGCGCA GTCTTGCTAGACGAAGCTTTTGATCTATATTGCCGGGGATTACTGATCGGATTTGGCCCG TTTTGGGAACACATGTTGGGATACTGGAGAGAGAGCTTGAAGAGGCCAGAGAAAGTCTTA TTTTTAAAGTACGAGGATCTCAAAGAAGACATCGAGACCAACTTGAAGAAGCTAGCAAGT TTCTTAGGACTTCCTTTCACCGAAGAAGAGGGAACAAAAGGGAGTTGTGAAAGCTATCGCT GATCTGTGTAGCTTTGAGAATCTGAAGAAGTTGGAGGTGAACAAGTCAAGCAAATTGATC $\tt TTGTCGCCATCACAAGTGGAAAGATTGTCAGCCTTAGTGGATGACAAGTTAGCTGGATCT$ GGTCTCACTTTCAGATTGAGT

Figure 10

SEQ ID NO 4:

MAIPSFSMCHKPELLKEGKSEGQEEEGLSYEFQEMLDSLPKERGRRNRYLYLFQGFRCQA KEIQAITSFQKHFQSLPDDVVLATIPKSGTTWLKALTFTILTRHRFDPVSSSSSDHPLLT SNPHDLVPFFEYKLYANGNVPDLSGLASPRTFATHVPFGALKDSVENPSVKVVYLCRNPF DTFISMWHYINNITSESVSAVLLDEAFDLYCRGLLIGFGPFWEHMLGYWRESLKRPEKVL FLKYEDLKEDIETNLKKLASFLGLPFTEEEEQKGVVKAIADLCSFENLKKLEVNKSSKLI QNYENRFLFRKGEVSDLVNYLSPSQVERLSALVDDKLAGSGLTFRLS

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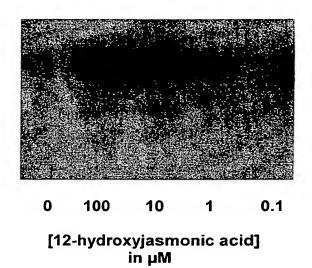


Fig. 11

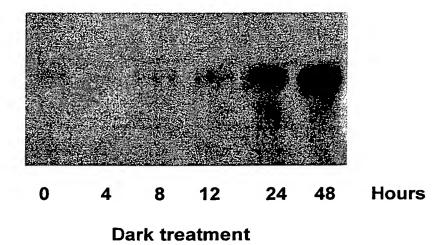


Fig. 12